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## SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (i) APPLICANT: Frank, Bruce L.
    Roberts, Peter C.
    Goodchild, John
    Craig, J. Charles
    Mills, John S.
    Slade, Andrew
    Roberts, Noel A.
    Jupp, Raymond
  - (ii) TITLE OF INVENTION: OLIGONUCLEOTIDES SPECIFIC FOR HEPATITIS B VIRUS
  - (iii) NUMBER OF SEQUENCES: 56
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: Lappin & Kusmer
    - (B) STREET: 200 State Street
    - (C) CITY: Boston
    - (D) STATE: MA
    - (E) COUNTRY: USA
    - (F) ZIP: 02109
    - (v) COMPUTER READABLE FORM:
      - (A) MEDIUM TYPE: Floppy disk
      - (B) COMPUTER: IBM PC compatible
      - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
  - (vi) CURRENT APPLICATION DATA:
    - (A) APPLICATION NUMBER: US
    - (B) FILING DATE:
    - (C) CLASSIFICATION:
  - (viii) ATTORNEY/AGENT INFORMATION:
    - (A) NAME: Kerner, Ann-Louise
    - (B) REGISTRATION NUMBER: 33,523
    - (C) REFERENCE/DOCKET NUMBER: HYZ-041
    - (ix) TELECOMMUNICATION INFORMATION:
      - (A) TELEPHONE: 617-330-1300
      - (B) TELEFAX: 617-330-1311
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:

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	(A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GGTGCGCA	GA CCAATTTATG	20
(2) INFO	RMATION FOR SEQ ID NO:2:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CATGGTGC	TG GTGCGCAGA	19
(2) INFO	RMATION FOR SEQ ID NO:3:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GAAAAAGT	TG CATGGTGCTG	20

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(2) INFO	RMATION FOR SEQ ID NO:4:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GAGGTGAA	AA AGTTGCATGG	20
(2) INFO	RMATION FOR SEQ ID NO:5:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
AGGCAGAG	GT GAAAAGTTG	20
(2) INFO	RMATION FOR SEQ ID NO:6:	
(1)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 12 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

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AGGCAGAG	FT GA	12
(2) INFO	RMATION FOR SEQ ID NO:7:	
( <b>i</b> )	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
AGAGATGA:	TT AGGCAGAGGT	20
(2) INFO	RMATION FOR SEQ ID NO:8:	
(1)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
GACATGAA	CA AGAGATGATT AGGCAGAGGT	30
(2) INFO	RMATION FOR SEQ ID NO:9:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	•
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	

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(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	
GACATGTAC	AGAGATGATT	20
(2) INFOR	RMATION FOR SEQ ID NO:10:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
GTAGGACA!	IG AACAAGAGAT	20
(2) INFO	RMATION FOR SEQ ID NO:11:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	
TTGGAGGC'	TT GAACAGTAGG	20
(2) INFO	RMATION FOR SEQ ID NO:12:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	

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(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CACAGCTTC	G AGGCTTGAAC	20
(2) INFO	RMATION FOR SEQ ID NO:13:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:13:	
AGCCACCC	AA GGCACAGCTT	20
(2) INFO	RMATION FOR SEQ ID NO:14:	
<b>(i)</b>	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:14:	
TCGATGTC	CA TGCCCCAAAG	20
(2) INFO	RMATION FOR SEQ ID NO:15:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	ΜΟΙ.Ε. ΤΥΡΕ - ΝΝΔ	

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(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:15:	
TAAGGGTC	GA TGTCCATGCC	20
(2) INFO	RMATION FOR SEQ ID NO:16:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:16:	
TTATAAGG	GT CGATGTCCAT	20
(2) INFO	RMATION FOR SEQ ID NO:17:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	•
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:17:	
AAATTCTT	TA TAAGGGTCGA TGTCCAT	27
(2) INFO	RMATION FOR SEQ ID NO:18:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:18:	
TATAAGGG	TC GA	12
(2) INFO	RMATION FOR SEQ ID NO:19:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	•
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:19:	
AAATTCTT	TA TAAGGGTCGA	20
(2) INFO	RMATION FOR SEQ ID NO:20:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:20:	
GTATCTAG	BAA GATCTCGTAC	20
(2) INFO	DRMATION FOR SEQ ID NO:21:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid	

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	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) 1	MOLECULE TYPE: DNA	
(iii) 1	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:21:	
GCGGTGTCT	'A GAAGATCTCG	20
(2) INFOR	MATION FOR SEQ ID NO:22:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA .	
(iii)	HYPOTHETICAL: NO	-
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:22:	
GAGGCGGTG	T CTAGGAGATC	20
(2) INFOR	RMATION FOR SEQ ID NO:23:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:23:	
GAGCTGAGG	GC GGTGTCTAGG	20
(2) INFOR	RMATION FOR SEQ ID NO:24:	

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(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:24:	
ATACAGAG	CT GAGGCGGTAT	20
(2) INFO	RMATION FOR SEQ ID NO:25:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:25:	
TCCCGATA	CA GAGCTGAGGC	20
(2) INFO	RMATION FOR SEQ ID NO:26:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:26:	
AGGCTTCC	CG ATACAGAGCT	20
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(2) INFOR	MATION FOR SEQ ID NO:27:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:27:	
ACAATGCT	CA GGAGACTCTA	20
(2) INFO	RMATION FOR SEQ ID NO:28:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:28:	
GCAGTATG	GAGGTGAGCA	20
(2) INFO	RMATION FOR SEQ ID NO:29:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEOUENCE DESCRIPTION: SEO ID NO:29:	

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GAGTGCAGTA TGGTGAGGTG

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(2) TNEO	RMATION FOR SEQ ID NO:30:	
	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:30:	
TGCCTGAG	TG CAGTATGGTG	20
(2) INFO	RMATION FOR SEQ ID NO:31:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	;
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(vi)	ANTI-SENSE: YES	,
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:31:	
TTGCTTGC	CT GAGTGCAGTA	20
(2) INFO	RMATION FOR SEQ ID NO:32:	
<b>(i)</b>	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	

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(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:32:	
TAAGGGTC	BA AGAGATGATT	20
(2) INFO	RMATION FOR SEQ ID NO:33:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:33:	
AGAGATGA'	TT TAAGGGTCGA	20
(2) INFO	RMATION FOR SEQ ID NO:34:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:34:	
TAAGGGTC	GA AGGCAGAGGT	20
(2) INFO	RMATION FOR SEQ ID NO:35:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	

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(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:35:		
AGGCAGAG	GT TAAGGGTCGA		20
(2) INFO	RMATION FOR SEQ ID NO:36:		
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:36:		
TATAAGGG	STC GAAGGCAGAG GTGA		24
(2) INFO	DRMATION FOR SEQ ID NO:37:		
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	: •	
(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:37:		
AGGCAGAG	GGT GATATAAGGG TCGA		24
(2) INFO	ORMATION FOR SEQ ID NO:38:		
(i)	) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
(ii)	) MOLECULE TYPE: DNA		

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(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:38:	
AGAGATGA!	TT AGGCAGAGGT TAAGGGTCGA	30
(2) INFO	RMATION FOR SEQ ID NO:39:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ 1D NO:39:	
GACATGAA	CA AGAGATGATT TAAGGGTCGA	30
(2) INFO	RMATION FOR SEQ ID NO:40:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:40:	
AGAGATGA	TT TAAGGGTCGA TGTCCATGCC	30
(2) INFO	RMATION FOR SEQ ID NO:41:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:	
i	AGGCAGAG	GT TAAGGGTCGA TGTCCATGCC	. 30
	(2) INFO	RMATION FOR SEQ ID NO:42:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:	
	GGCATTTG	GT GGTCTATAAG	20
	(2) INFO	RMATION FOR SEQ ID NO:43:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:	
	GATAGGGG	GCA TTTGGTGGTC	20
	(2) INFO	ORMATION FOR SEQ ID NO:44:	
	(1)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid	

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	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:44;	
TGTTGATA	GG ATAGGGGCAT	20
(2) INFO	RMATION FOR SEQ ID NO:45:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	÷
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:45:	
ACCCAAGG	CA CAGCTTGGAG	20
(2) INFO	RMATION FOR SEQ ID NO:46:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:46:	
GACAGGGG	CA TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:47:	

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(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:47:	
GATAGGGG	CC TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:48:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
( <b>ii</b> )	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:48:	
GATAGGGG	CA TTTGGTGCTC	20
(2) INFO	RMATION FOR SEQ ID NO:49:	
<b>(i)</b>	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:49:	
GACAGGGG	CC TTTGGTGCTC	20

(2) INFORMATION FOR SEQ ID NO:50:

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(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(i.i)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:50:	
ATGTTTTT	GG CGTCTTCCAT	20
(2) INFO	RMATION FOR SEQ ID NO:51:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:51:	
TGCTGGGG	GG AACTAATGAC T	21
(2) INFO	RMATION FOR SEQ ID NO:52:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:52:	
	•	

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GGAGTGCGAA TCCACACTCC GAAAG

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(2) INFO	RMATION FOR SEQ ID NO:53:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:53:	
TAAGGGTC	GA	10
(2) INFO	RMATION FOR SEQ ID NO:54:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
· (ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:54:	
AGGCAGAG	GT	10
(2) INFO	RMATION FOR SEQ ID NO:55:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:	
ATATTCGCAC GAATTCAGCT TATAGACCAC CAAATG	36
(2) INFORMATION FOR SEQ ID NO:56:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 34 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:	
GTTGGGATTG CTGCAGCTAT CTGGATTTGC GGTG	34